

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/553,661
Source: PT/10
Date Processed by STIC: 10/28/05

ENTERED

**Raw Sequence Listing before editing
(for reference only)**



PCT

RAW SEQUENCE LISTING

DATE: 10/28/2005

PATENT APPLICATION: US/10/553,661

TIME: 10:22:25

Input Set : A:\50680-6 Seq 07-10-05 v2.txt

Output Set: N:\CRF4\10282005\J553661.raw

3 <110> APPLICANT: Genesis Group Inc., Kenneth Kao, Catherine Popadiuk
5 <120> TITLE OF INVENTION: Pygopus in Diagnosis and Treatment of Cancer
7 <130> FILE REFERENCE: 50680-4
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/553,661
C--> 9 <141> CURRENT FILING DATE: 2005-10-17
9 <150> PRIOR APPLICATION NUMBER: US 60/463 309
10 <151> PRIOR FILING DATE: 2003-04-17
12 <150> PRIOR APPLICATION NUMBER: US 60/496 012
13 <151> PRIOR FILING DATE: 2003-08-19
15 <160> NUMBER OF SEQ ID NOS: 28
17 <170> SOFTWARE: PatentIn version 3.3

ERRORED SEQUENCES

757 <210> SEQ ID NO: 28
758 <211> LENGTH: 19
759 <212> TYPE: DNA
760 <213> ORGANISM: artificial
762 <220> FEATURE:
763 <223> OTHER INFORMATION: mismatched sequence to Hpy5
765 <400> SEQUENCE: 28
766 gtctgagcta atcattggt 19
E--> 767 1
E--> 769 1/14

Does Not Comply
Corrected Diskette Needed

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 10/28/2005
PATENT APPLICATION: US/10/553,661 TIME: 10:22:26

Input Set : A:\50680-6 Seq 07-10-05 v2.txt
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Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22,23,24,25,26,27,28

VERIFICATION SUMMARY

DATE: 10/28/2005

PATENT APPLICATION: US/10/553,661

TIME: 10:22:26

Input Set : A:\50680-6 Seq 07-10-05 v2.txt

Output Set: N:\CRF4\10282005\J553661.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application No
L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:767 M:254 E: No. of Bases conflict, this line has no nucleotides.
L:769 M:254 E: No. of Bases conflict, LENGTH:Input:14 Counted:20 SEQ:28
L:769 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:769 M:252 E: No. of Seq. differs, <211> LENGTH:Input:19 Found:20 SEQ:28